

#23

Patent  
Attorney's Docket No. 030560-057

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

RECEIVED  
JUL 16 2003  
TECH CENTER 1600/2900

In re Patent Application of	)	
	)	
Friedrich ALTMANN	)	Group Art Unit: 1635
	)	
Serial No.: 09/913,858	)	Examiner: Sean McGarry
	)	
Filed: August 20, 2001	)	ATTENTION: BOX SEQUENCE
	)	
For: FUCOSYL TRANSFERASE GENE	)	

**DECLARATION PURSUANT TO**  
**37 C.F.R. §§1.821-1.825**

Commissioner for Patents  
P.O. Box 1450  
Alexandria, VA 22313-1450

Sir:

I, Donna M. Meuth, declare as follows:

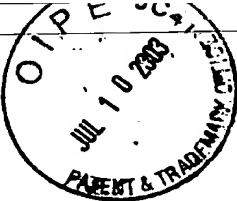
1. That the content of the paper and computer readable copies of the Sequence Listing, submitted in accordance with 37 C.F.R. §1.821(c) and (e), respectively, are the same in compliance with §1.821(f).
2. That the submission, filed in accordance with 37 C.F.R. §1.821(g)[or (h)], herein does not include new matter [or go beyond the disclosure in the international application].

Serial No.: 09/913,858

I hereby declare that all statements made herein of my own knowledge are true and that all statements were made on information and belief and are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

Date July 10, 2003

Donna M. Meuth  
Donna M. Meuth  
Registration No. 36,607



1635

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For: FUCOSYL TRANSFERASE GENE	)	

TRANSMITTAL LETTER

Commissioner for Patents  
P.O. Box 1450  
Alexandria, VA 22313-1450

Sir:

In complete response to the Office communication dated June 11, 2003,  
enclosed please find:

- [X] A copy of the "Sequence Listing" in computer readable form in compliance with 37 C.F.R. §§1.823(b) and 1.824.
- [X] A statement that the content of the paper and computer readable copies are the same as set forth in 37 C.F.R. §1.821(f).

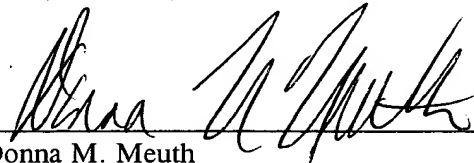
The Commissioner is hereby authorized to charge any additional fees under 37 C.F.R. §§1.16, 1.17, and 1.21 that may be required by this paper, and to credit any overpayment to Deposit Account No. 02-4800. A duplicate copy of this paper is enclosed.

Respectfully submitted,

P.O. Box 1404  
Alexandria, VA 22313-1404  
(703) 836-6620

BURNS, DOANE, SWECKER & MATHIS, L.L.P.

Date: July 16, 2003

By   
Donna M. Meuth  
Registration No. 36,607

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1600

## RAW SEQUENCE LISTING

DATE: 07/15/2003

PATENT APPLICATION: US/09/913,858B

TIME: 10:50:53

Input Set : A:\030560-057.ST25.txt

Output Set: N:\CRF4\07152003\I913858B.raw

4 <110> APPLICANT: Altmann, Friedrich  
6 <120> TITLE OF INVENTION: Fucosyl Transferase Gene  
8 <130> FILE REFERENCE: 030560-057  
10 <140> CURRENT APPLICATION NUMBER: US 09/913,858B  
11 <141> CURRENT FILING DATE: 2001-08-20  
13 <150> PRIOR APPLICATION NUMBER: PCT/AT00/00040  
14 <151> PRIOR FILING DATE: 2000-02-17  
16 <150> PRIOR APPLICATION NUMBER: AT A 270/99  
17 <151> PRIOR FILING DATE: 1999-02-18  
19 <160> NUMBER OF SEQ ID NOS: 31  
21 <170> SOFTWARE: PatentIn version 3.1  
23 <210> SEQ ID NO: 1  
25 <211> LENGTH: 2198  
26 <212> TYPE: DNA  
27 <213> ORGANISM: Unknown Organism  
29 <220> FEATURE:  
30 <223> OTHER INFORMATION: Description of Unknown Organism: plant  
32 <400> SEQUENCE: 1

ENTERED

33 actaactcaa acgctgcatt ttcttttttc ttccaggaa ccatccaccc ataacaacaa 60  
34 aaaaaacaac agcaagctgt gtttttttta tcgttctttt tctttaaaca agcaccacca 120  
35 tcatggaatc gtgctcataa cgccaaaatt ttccatttcc ctttgatttt tagtttattt 180  
36 tgcggaattg gcagttgggg gcgcaattga atgatgggtc tgttgacgaa tcttcgaggc 240  
37 tcgagaacag atggtgcccc acaagacagc ttaccggtt tggctccggg aggcaaccca 300  
38 aagaggaaat ggagcaatct aatgcctctt gttgttgccc ttgtggtcat cgcgagatc 360  
39 gcgtttctgg gtaggttgga tatggccaaa aacgccgcca tgggtgactc cctcgtgac 420  
40 ttcttctacc gctctcgagc ggtcgttgaa ggtgacgatt tgggggttggg tttggtggct 480  
41 tctgatcgga attctgaatc gtatagttgt gaggaatggt tggagaggga ggatgctgtc 540  
42 acgtattcga ggggcttttc caaagagcct atttttgttt ctggagctga tcaggagtgg 600  
43 aagtcgtggt cggttggtatg taaatttggg tttagtgggg atagaaagcc agatgccgca 660  
44 tttgggttac ctcaaccaag tggaacagct agcattctgc gatcaatgga atcagcagaa 720  
45 tactatgctg agaacaatat tgccatggca agacggaggg gatataacat cgtaatgaca 780  
46 accagtctat cttcggatgt tcctgttgga tatttttcat gggctgagta tgatatgatg 840  
47 gcaccagtgc agccgaaaac tgaagctgct cttgcagctg ctttcatttc caattgtggt 900  
48 gctcgaaatt tccggttgca agctcttgag gcccttgaaa aatcaaacat caaaattgat 960  
49 tcttatggtg gttgtcacag gaaccgtgat ggaagagtga acaaagtga agccctgaag 1020  
50 cactacaaat ttagcttagc gtttgaaaat tcgaatgagg aagattatgt aactgaaaaa 1080  
51 ttcttccaat cccttgttgc tggaactgtc cctgtggttg ttggtgctcc aaatattcag 1140  
52 gactttgctc cttctcctgg ttcaatttta catattaaag agatagagga tgttgagtct 1200  
53 gttgcaaaga ccatgagata tctagcagaa aaatcccgaag catataatca atcattgagg 1260  
54 tggaagtatg aggggtccatc tgactccttc aaggcccttg tggatatggc agctgtgcat 1320  
55 tcatcggtgc gtctttgcat tcaacttgcc acagttagta gagagaagga agaaaataat 1380  
56 ccaagcctta agagacgtcc ttgcaagtgc actagagggc cagaaaccgt atatcatatc 1440  
57 tatgtcagag aaaggggaag gtttgagatg gagtccattt acctgaggtc tagcaattta 1500

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Output Set: N:\CRF4\07152003\I913858B.raw

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58 actctgaatg ctgtgaaggc tgctgttggt ttgaagttca catccctgaa tcttgtgcct 1560
59 gtatggaaga ctgaaaggcc tgaagttata agagggggga gtgctttaa actctacaaa 1620
60 atatacccaa ttggcttgac acagagacaa gctctttata ccttcagctt caaaggtgat 1680
61 gctgatttca ggagtcactt ggagaacaat ccttgtgcc aagttgaagt catttttgtg 1740
62 tagcatgccc taaatggtac ctctgctcta cctgaattag cttcacttag ctgagcacta 1800
63 gctagagttt taggaatgag tatggcagtg aatatggcat ggctttattt atgcctagtt 1860
64 tcttggccaa ctcatgtatg ttttgtataa gacatcacac ttttaattta aacttgtttc 1920
65 tgtagaagtg caaatccata tttaatgctt agtttttagtg ctcttatctg atcatctaga 1980
66 agtcacagtt cttgtatatt gtgagtgaat actgaaatct aatagaagga tcagatgttt 2040
67 cactcaagac acattattac ttcattgttg tttgatgac tcgagctttt ttagtgtctg 2100
68 gaactgtccc tgtggtttga gcaectgtta ttgcttcagt gttactgtcc agtgggtatc 2160
69 gtttttgacc tctaaaaaaa aaaaaaaaaa aaaaaaaaaa 2198

```

71 &lt;210&gt; SEQ ID NO: 2

72 &lt;211&gt; LENGTH: 510

73 &lt;212&gt; TYPE: PRT

74 &lt;213&gt; ORGANISM: Unknown Organism

76 &lt;220&gt; FEATURE:

77 &lt;223&gt; OTHER INFORMATION: Description of Unknown Organism:plant

79 &lt;400&gt; SEQUENCE: 2

```

80 Met Met Gly Leu Leu Thr Asn Leu Arg Gly Ser Arg Thr Asp Gly Ala
81 1 5 10 15
83 Gln Gln Asp Ser Leu Pro Val Leu Ala Pro Gly Gly Asn Pro Lys Arg
84 20 25 30
86 Lys Trp Ser Asn Leu Met Pro Leu Val Val Ala Leu Val Val Ile Ala
87 35 40 45
89 Glu Ile Ala Phe Leu Gly Arg Leu Asp Met Ala Lys Asn Ala Ala Met
90 50 55 60
92 Val Asp Ser Leu Ala Asp Phe Phe Tyr Arg Ser Arg Ala Val Val Glu
93 65 70 75 80
95 Gly Asp Asp Leu Gly Leu Gly Leu Val Ala Ser Asp Arg Asn Ser Glu
96 85 90 95
98 Ser Tyr Ser Cys Glu Glu Trp Leu Glu Arg Glu Asp Ala Val Thr Tyr
99 100 105 110
101 Ser Arg Gly Phe Ser Lys Glu Pro Ile Phe Val Ser Gly Ala Asp Gln
102 115 120 125
104 Glu Trp Lys Ser Cys Ser Val Gly Cys Lys Phe Gly Phe Ser Gly Asp
105 130 135 140
107 Arg Lys Pro Asp Ala Ala Phe Gly Leu Pro Gln Pro Ser Gly Thr Ala
108 145 150 155 160
110 Ser Ile Leu Arg Ser Met Glu Ser Ala Glu Tyr Tyr Ala Glu Asn Asn
111 165 170 175
113 Ile Ala Met Ala Arg Arg Arg Gly Tyr Asn Ile Val Met Thr Thr Ser
114 180 185 190
116 Leu Ser Ser Asp Val Pro Val Gly Tyr Phe Ser Trp Ala Glu Tyr Asp
117 195 200 205
119 Met Met Ala Pro Val Gln Pro Lys Thr Glu Ala Ala Leu Ala Ala Ala
120 210 215 220
122 Phe Ile Ser Asn Cys Gly Ala Arg Asn Phe Arg Leu Gln Ala Leu Glu
123 225 230 235 240

```

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TIME: 10:50:53

Input Set : A:\030560-057.ST25.txt

Output Set: N:\CRF4\07152003\I913858B.raw

```

125 Ala Leu Glu Lys Ser Asn Ile Lys Ile Asp Ser Tyr Gly Gly Cys His
126                245                250                255
128 Arg Asn Arg Asp Gly Arg Val Asn Lys Val Glu Ala Leu Lys His Tyr
129                260                265                270
131 Lys Phe Ser Leu Ala Phe Glu Asn Ser Asn Glu Glu Asp Tyr Val Thr
132                275                280                285
134 Glu Lys Phe Phe Gln Ser Leu Val Ala Gly Thr Val Pro Val Val Val
135                290                295                300
137 Gly Ala Pro Asn Ile Gln Asp Phe Ala Pro Ser Pro Gly Ser Ile Leu
138 305                310                315                320
140 His Ile Lys Glu Ile Glu Asp Val Glu Ser Val Ala Lys Thr Met Arg
141                325                330                335
143 Tyr Leu Ala Glu Asn Pro Glu Ala Tyr Asn Gln Ser Leu Arg Trp Lys
144                340                345                350
146 Tyr Glu Gly Pro Ser Asp Ser Phe Lys Ala Leu Val Asp Met Ala Ala
147                355                360                365
149 Val His Ser Ser Cys Arg Leu Cys Ile His Leu Ala Thr Val Ser Arg
150                370                375                380
152 Glu Lys Glu Glu Asn Asn Pro Ser Leu Lys Arg Arg Pro Cys Lys Cys
153 385                390                395                400
155 Thr Arg Gly Pro Glu Thr Val Tyr His Ile Tyr Val Arg Glu Arg Gly
156                405                410                415
158 Arg Phe Glu Met Glu Ser Ile Tyr Leu Arg Ser Ser Asn Leu Thr Leu
159                420                425                430
161 Asn Ala Val Lys Ala Ala Val Val Leu Lys Phe Thr Ser Leu Asn Leu
162                435                440                445
164 Val Pro Val Trp Lys Thr Glu Arg Pro Glu Val Ile Arg Gly Gly Ser
165                450                455                460
167 Ala Leu Lys Leu Tyr Lys Ile Tyr Pro Ile Gly Leu Thr Gln Arg Gln
168 465                470                475                480
170 Ala Leu Tyr Thr Phe Ser Phe Lys Gly Asp Ala Asp Phe Arg Ser His
171                485                490                495
173 Leu Glu Asn Asn Pro Cys Ala Lys Phe Glu Val Ile Phe Val
174                500                505                510

```

177 &lt;210&gt; SEQ ID NO: 3

178 &lt;211&gt; LENGTH: 105

179 &lt;212&gt; TYPE: DNA

180 &lt;213&gt; ORGANISM: Artificial Sequence

182 &lt;220&gt; FEATURE:

183 &lt;223&gt; OTHER INFORMATION: Description of Artificial Sequence: GlcNAc-alpha1,3-fucosyl

185 &lt;400&gt; SEQUENCE: 3

186 gaagccctga agcactacaa atttagctta gcgtttgaaa attcgaatga ggaagattat 60

187 gtaactgaaa aattcttcca atcccttggt gctggaactg tccct 105

189 &lt;210&gt; SEQ ID NO: 4

190 &lt;211&gt; LENGTH: 35

191 &lt;212&gt; TYPE: PRT

192 &lt;213&gt; ORGANISM: Artificial Sequence

194 &lt;220&gt; FEATURE:

195 &lt;223&gt; OTHER INFORMATION: Description of Artificial Sequence: Mung bean

## RAW SEQUENCE LISTING

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```

197 <400> SEQUENCE: 4
198 Glu Ala Leu Lys His Tyr Lys Phe Ser Leu Ala Phe Glu Asn Ser Asn
199   1           5           10           15
201 Glu Glu Asp Tyr Val Thr Glu Lys Phe Phe Gln Ser Leu Val Ala Gly
202           20           25           30
204 Thr Val Pro
205           35
208 <210> SEQ ID NO: 5
209 <211> LENGTH: 15
210 <212> TYPE: PRT
211 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223> OTHER INFORMATION: Description of Artificial Sequence:n-terminal sequence
215     of tryptic peptide
217 <220> FEATURE:
218 <221> NAME/KEY: MISC_FEATURE
219 <222> LOCATION: (5)..(5)
220 <223> OTHER INFORMATION: Xaa = any amino acid
222 <400> SEQUENCE: 5
W--> 224 Lys Pro Asp Ala Xaa Phe Gly Leu Pro Gln Pro Ser Thr Ala Ser
225   1           5           10           15
230 <210> SEQ ID NO: 6
231 <211> LENGTH: 10
232 <212> TYPE: PRT
233 <213> ORGANISM: Artificial Sequence
235 <220> FEATURE:
236 <223> OTHER INFORMATION: Description of Artificial Sequence:n-terminal sequence
237     of tryptic peptide
239 <400> SEQUENCE: 6
240 Pro Glu Thr Val Tyr His Ile Tyr Val Arg
241   1           5           10
244 <210> SEQ ID NO: 7
245 <211> LENGTH: 13
246 <212> TYPE: PRT
247 <213> ORGANISM: Artificial Sequence
249 <220> FEATURE:
250 <223> OTHER INFORMATION: Description of Artificial Sequence:n-terminal sequence
251     of tryptic peptide
253 <400> SEQUENCE: 7
254 Met Glu Ser Ala Glu Tyr Tyr Ala Glu Asn Asn Ile Ala
255   1           5           10
258 <210> SEQ ID NO: 8
259 <211> LENGTH: 10
260 <212> TYPE: PRT
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence:n-terminal sequence
265     of tryptic peptide
267 <400> SEQUENCE: 8

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## RAW SEQUENCE LISTING

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Input Set : A:\030560-057.ST25.txt

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```

268 Gly Arg Phe Glu Met Glu Ser Ile Tyr Leu
269   1           5           10
271 <210> SEQ ID NO: 9
272 <211> LENGTH: 29
273 <212> TYPE: DNA
274 <213> ORGANISM: Artificial Sequence
276 <220> FEATURE:
277 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
279 <220> FEATURE:
280 <221> NAME/KEY: misc_feature
281 <222> LOCATION: (3)..(15)
282 <223> OTHER INFORMATION: n = any nucleotide
284 <400> SEQUENCE: 9
W--> 285 gngartayt aygngaraa yaayathgc 29
288 <210> SEQ ID NO: 10
289 <211> LENGTH: 22
290 <212> TYPE: DNA
291 <213> ORGANISM: Artificial Sequence
293 <220> FEATURE:
294 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
296 <220> FEATURE:
297 <221> NAME/KEY: misc_feature
298 <222> LOCATION: (14)..(17)
299 <223> OTHER INFORMATION: n = any nucleotide
301 <400> SEQUENCE: 10
W--> 302 crtadatrtg rtanacngty tc 22
305 <210> SEQ ID NO: 11
306 <211> LENGTH: 20
307 <212> TYPE: DNA
308 <213> ORGANISM: Artificial Sequence
310 <220> FEATURE:
311 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
313 <220> FEATURE:
314 <221> NAME/KEY: misc_feature
315 <222> LOCATION: (6)..(6)
316 <223> OTHER INFORMATION: n = any nucleotide
318 <400> SEQUENCE: 11
W--> 319 tadatnswyt ccatytcraa 20
322 <210> SEQ ID NO: 12
323 <211> LENGTH: 20
324 <212> TYPE: DNA
325 <213> ORGANISM: Artificial Sequence
327 <220> FEATURE:
328 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
330 <400> SEQUENCE: 12
331 ctggaactgt ccctgtggtt 20
333 <210> SEQ ID NO: 13
334 <211> LENGTH: 20
335 <212> TYPE: DNA

```



RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/913,858B

DATE: 07/15/2003  
TIME: 10:50:54

Input Set : A:\030560-057.ST25.txt  
Output Set: N:\CRF4\07152003\I913858B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 5  
Seq#:9; N Pos. 3,15  
Seq#:10; N Pos. 14,17  
Seq#:11; N Pos. 6

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**VERIFICATION SUMMARY**

DATE: 07/15/2003

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Input Set : A:\030560-057.ST25.txt

Output Set: N:\CRF4\07152003\I913858B.raw

L:224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
L:285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
L:302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
L:319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0

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